

SEQUENCE LISTING

<110> University of Utah Research Foundation  
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Li, Dean Y.

<120> ELASTIN-BASED COMPOSITIONS

<130> 22458-702

<140> Not Yet Assigned

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<150> PCT/US00/02526

<151> 2000-02-28

<150> US 09/258,217

<151> 1999-02-26

<160> 8

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 6

<212> PRT

<213> Homo sapiens

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<210> 2

<211> 42

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<213> Artificial Sequence

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<223> Synthetic

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20 25 30  
Val Ala Pro Gly Val Gly Val Ala Pro Gly  
35 40

<210> 3  
 <211> 712  
 <212> PRT  
 <213> Homo sapiens

<400> 3

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Leu	Leu	Ser	Ile	Leu	His	Pro	Ser	Arg	Pro	Gly	Gly	Val	Pro	Gly	Ala			
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Ile	Pro	Gly	Gly	Val	Pro	Gly	Gly	Val	Phe	Tyr	Pro	Gly	Ala	Gly	Leu			
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Gly	Ala	Leu	Gly	Gly	Gly	Ala	Leu	Gly	Pro	Gly	Gly	Lys	Pro	Leu	Lys			
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Pro	Val	Pro	Gly	Gly	Leu	Ala	Gly	Ala	Gly	Leu	Gly	Ala	Gly	Leu	Gly			
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Ala	Phe	Pro	Ala	Val	Thr	Phe	Pro	Gly	Ala	Leu	Val	Pro	Gly	Gly	Val			
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Ala	Asp	Ala	Ala	Ala	Ala	Tyr	Lys	Ala	Ala	Lys	Ala	Gly	Ala	Gly	Leu			
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Gly	Gly	Val	Pro	Gly	Val	Gly	Gly	Leu	Gly	Val	Ser	Ala	Gly	Ala	Val			
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Val	Pro	Gln	Pro	Gly	Ala	Gly	Val	Lys	Pro	Gly	Lys	Val	Pro	Gly	Val			
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Gly	Leu	Pro	Gly	Val	Tyr	Pro	Gly	Gly	Val	Leu	Pro	Gly	Ala	Arg	Phe			
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Pro	Gly	Val	Gly	Val	Leu	Pro	Gly	Val	Pro	Thr	Gly	Ala	Gly	Val	Lys			
				165					170						175			
Pro	Lys	Ala	Pro	Gly	Val	Gly	Gly	Ala	Phe	Ala	Gly	Ile	Pro	Gly	Val			
			180					185						190				
Gly	Pro	Phe	Gly	Gly	Pro	Gln	Pro	Gly	Val	Pro	Leu	Gly	Tyr	Pro	Ile			
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Lys	Ala	Pro	Lys	Leu	Pro	Gly	Gly	Tyr	Gly	Leu	Pro	Tyr	Thr	Thr	Gly			
	210					215					220							
Lys	Leu	Pro	Tyr	Gly	Tyr	Gly	Pro	Gly	Gly	Val	Ala	Gly	Ala	Ala	Gly			
225					230					235					240			
Lys	Ala	Gly	Tyr	Pro	Thr	Gly	Thr	Gly	Val	Gly	Pro	Gln	Ala	Ala	Ala			
				245					250					255				
Ala	Ala	Ala	Ala	Lys	Ala	Ala	Ala	Lys	Phe	Gly	Ala	Gly	Ala	Ala	Gly			
			260					265					270					
Val	Leu	Pro	Gly	Val	Gly	Gly	Ala	Gly	Val	Pro	Gly	Val	Pro	Gly	Ala			
		275					280					285						
Ile	Pro	Gly	Ile	Gly	Gly	Ile	Ala	Gly	Val	Gly	Thr	Pro	Ala	Ala	Ala			
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Ala	Ala	Ala	Ala	Ala	Ala	Ala	Lys	Ala	Ala	Lys	Tyr	Gly	Ala	Ala	Ala			
305					310					315					320			
Gly	Leu	Val	Pro	Gly	Gly	Pro	Gly	Phe	Gly	Pro	Gly	Val	Val	Gly	Val			
				325					330					335				
Pro	Gly	Ala	Gly	Val	Pro	Gly	Val	Gly	Val	Pro	Gly	Ala	Gly	Ile	Pro			
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Ser	Pro	Glu	Ala	Ala	Ala	Lys	Ala	Ala	Ala	Lys	Ala	Ala	Lys	Tyr	Gly
	370					375					380				
Ala	Arg	Pro	Gly	Val	Gly	Val	Gly	Gly	Ile	Pro	Thr	Tyr	Gly	Val	Gly
385					390					395					400
Ala	Gly	Gly	Phe	Pro	Gly	Phe	Gly	Val	Gly	Val	Gly	Gly	Ile	Pro	Gly
				405					410					415	
Val	Ala	Gly	Val	Pro	Gly	Val	Gly	Gly	Val	Pro	Gly	Val	Gly	Gly	Val
			420					425					430		
Pro	Gly	Val	Gly	Ile	Ser	Pro	Glu	Ala	Gln	Ala	Ala	Ala	Ala	Ala	Lys
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Ala	Ala	Lys	Tyr	Gly	Val	Gly	Thr	Pro	Ala	Ala	Ala	Ala	Ala	Lys	Ala
	450					455					460				
Ala	Ala	Lys	Ala	Ala	Gln	Phe	Ala	Leu	Leu	Asn	Leu	Ala	Gly	Leu	Val
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Val	Ala	Pro	Gly	Val	Gly	Leu	Ala	Pro	Gly	Val	Gly	Val	Ala	Pro	Gly
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Val	Gly	Val	Ala	Pro	Gly	Val	Gly	Val	Ala	Pro	Gly	Ile	Gly	Pro	Gly
		515					520					525			
Gly	Val	Ala	Ala	Ala	Ala	Lys	Ser	Ala	Ala	Lys	Val	Ala	Ala	Lys	Ala
	530					535					540				
Gln	Leu	Arg	Ala	Ala	Ala	Gly	Leu	Gly	Ala	Gly	Ile	Pro	Gly	Leu	Gly
545					550					555					560
Val	Gly	Val	Gly	Val	Pro	Gly	Leu	Gly	Val	Gly	Ala	Gly	Val	Pro	Gly
				565					570					575	
Leu	Gly	Val	Gly	Ala	Gly	Val	Pro	Gly	Phe	Gly	Ala	Val	Pro	Gly	Ala
			580					585					590		
Leu	Ala	Ala	Ala	Lys	Ala	Ala	Lys	Tyr	Gly	Ala	Ala	Val	Pro	Gly	Val
		595					600					605			
Leu	Gly	Gly	Leu	Gly	Ala	Leu	Gly	Gly	Val	Gly	Ile	Pro	Gly	Gly	Val
	610					615					620				
Val	Gly	Ala	Gly	Pro	Ala	Ala	Ala	Ala	Ala	Ala	Ala	Lys	Ala	Ala	Ala
625					630					635					640
Lys	Ala	Ala	Gln	Phe	Gly	Leu	Val	Gly	Ala	Ala	Gly	Leu	Gly	Gly	Leu
				645					650					655	
Gly	Val	Gly	Gly	Leu	Gly	Val	Pro	Gly	Val	Gly	Gly	Leu	Gly	Gly	Ile
			660					665					670		
Pro	Pro	Ala	Ala	Ala	Ala	Lys	Ala	Ala	Lys	Tyr	Gly	Val	Ala	Ala	Arg
		675					680					685			
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Lys</															

<210>	4
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<212>	PRT

<213> Artificial Sequence

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<223> Sequence encoded by forward primer

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<210> 5

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Forward primer

<400> 5

ctgctgctgc atatggcggg tctgacggcg

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<212> PRT

<213> Artificial Sequence

<220>

<223> Sequence encoded by complement to reverse primer

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Ala Cys Gly Arg Lys Arg Lys Gln Lys Leu Ile Ser Glu Glu Asp Leu  
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<210> 7

<211> 66

<212> DNA

<213> Artificial Sequence

<220>

<223> Reverse primer

<400> 7

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60

66

<210> 8

<211> 730

<212> PRT

<213> Artificial Sequence

<220>

&lt;223&gt; Human elastin-c-myc fusion

&lt;400&gt; 8

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			20					25						30		
Ser	Arg	Pro	Gly	Gly	Val	Pro	Gly	Ala	Ile	Pro	Gly	Gly	Val	Pro	Gly	
		35					40					45				
Gly	Val	Phe	Tyr	Pro	Gly	Ala	Gly	Leu	Gly	Ala	Leu	Gly	Gly	Gly	Ala	
	50					55					60					
Leu	Gly	Pro	Gly	Gly	Lys	Pro	Leu	Lys	Pro	Val	Pro	Gly	Gly	Leu	Ala	
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Gly	Ala	Gly	Leu	Gly	Ala	Gly	Leu	Gly	Ala	Phe	Pro	Ala	Val	Thr	Phe	
				85					90					95		
Pro	Gly	Ala	Leu	Val	Pro	Gly	Gly	Val	Ala	Asp	Ala	Ala	Ala	Ala	Tyr	
			100					105					110			
Lys	Ala	Ala	Lys	Ala	Gly	Ala	Gly	Leu	Gly	Gly	Val	Pro	Gly	Val	Gly	
		115					120						125			
Gly	Leu	Gly	Val	Ser	Ala	Gly	Ala	Val	Val	Pro	Gln	Pro	Gly	Ala	Gly	
	130					135					140					
Val	Lys	Pro	Gly	Lys	Val	Pro	Gly	Val	Gly	Leu	Pro	Gly	Val	Tyr	Pro	
145					150					155					160	
Gly	Gly	Val	Leu	Pro	Gly	Ala	Arg	Phe	Pro	Gly	Val	Gly	Val	Leu	Pro	
				165					170					175		
Gly	Val	Pro	Thr	Gly	Ala	Gly	Val	Lys	Pro	Lys	Ala	Pro	Gly	Val	Gly	
			180					185					190			
Gly	Ala	Phe	Ala	Gly	Ile	Pro	Gly	Val	Gly	Pro	Phe	Gly	Gly	Pro	Gln	
		195					200					205				
Pro	Gly	Val	Pro	Leu	Gly	Tyr	Pro	Ile	Lys	Ala	Pro	Lys	Leu	Pro	Gly	
	210					215					220					
Gly	Tyr	Gly	Leu	Pro	Tyr	Thr	Thr	Gly	Lys	Leu	Pro	Tyr	Gly	Tyr	Gly	
225					230					235					240	
Pro	Gly	Gly	Val	Ala	Gly	Ala	Ala	Gly	Lys	Ala	Gly	Tyr	Pro	Thr	Gly	
				245					250					255		
Thr	Gly	Val	Gly	Pro	Gln	Ala	Ala	Ala	Ala	Ala	Ala	Ala	Lys	Ala	Ala	
			260					265					270			
Ala	Lys	Phe	Gly	Ala	Gly	Ala	Ala	Gly	Val	Leu	Pro	Gly	Val	Gly	Gly	
		275					280					285				
Ala	Gly	Val	Pro	Gly	Val	Pro	Gly	Ala	Ile	Pro	Gly	Ile	Gly	Gly	Ile	
	290					295					300					
Ala	Gly	Val	Gly	Thr	Pro	Ala	Ala	Ala	Ala	Ala	Ala	Ala	Ala	Ala	Ala	
305					310						315				320	
Lys	Ala	Ala	Lys	Tyr	Gly	Ala	Ala	Ala	Gly	Leu	Val	Pro	Gly	Gly	Pro	
				325					330					335		
Gly	Phe	Gly	Pro	Gly	Val	Val	Gly	Val	Pro	Gly	Ala	Gly	Val	Pro	Gly	
			340					345					350			
Val	Gly	Val	Pro	Gly	Ala	Gly	Ile	Pro	Val	Val	Pro	Gly	Ala	Gly	Ile	
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Pro	Gly	Ala	Ala	Val	Pro	Gly	Val	Val	Ser	Pro	Glu	Ala	Ala	Ala	Lys	

370	375	380
Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Arg Pro Gly Val Gly Val		
385	390	395
Gly Gly Ile Pro Thr Tyr Gly Val Gly Ala Gly Gly Phe Pro Gly Phe		400
	405	410
Gly Val Gly Val Gly Gly Ile Pro Gly Val Ala Gly Val Pro Gly Val		415
	420	425
Gly Gly Val Pro Gly Val Gly Gly Val Pro Gly Val Gly Ile Ser Pro		430
	435	440
Glu Ala Gln Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Val Gly		445
450	455	460
Thr Pro Ala Ala Ala Ala Lys Ala Ala Lys Ala Ala Gln Phe		465
465	470	475
Ala Leu Leu Asn Leu Ala Gly Leu Val Pro Gly Val Gly Val Ala Pro		480
	485	490
Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Leu		495
	500	505
Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val		510
	515	520
Gly Val Ala Pro Gly Ile Gly Pro Gly Gly Val Ala Ala Ala Ala Lys		525
530	535	540
Ser Ala Ala Lys Val Ala Ala Lys Ala Gln Leu Arg Ala Ala Ala Gly		545
545	550	555
Leu Gly Ala Gly Ile Pro Gly Leu Gly Val Gly Val Gly Val Pro Gly		560
	565	570
Leu Gly Val Gly Ala Gly Val Pro Gly Leu Gly Val Gly Ala Gly Val		575
	580	585
Pro Gly Phe Gly Ala Val Pro Gly Ala Leu Ala Ala Ala Lys Ala Ala		590
	595	600
Lys Tyr Gly Ala Ala Val Pro Gly Val Leu Gly Gly Leu Gly Ala Leu		605
610	615	620
Gly Gly Val Gly Ile Pro Gly Gly Val Val Gly Ala Gly Pro Ala Ala		625
625	630	635
Ala Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu		640
	645	650
Val Gly Ala Ala Gly Leu Gly Gly Leu Gly Val Gly Gly Leu Gly Val		655
	660	665
Pro Gly Val Gly Gly Leu Gly Gly Ile Pro Pro Ala Ala Ala Lys		670
	675	680
Ala Ala Lys Tyr Gly Val Ala Ala Arg Pro Gly Phe Gly Leu Ser Pro		685
690	695	700
Ile Phe Pro Gly Gly Ala Cys Leu Gly Lys Ala Cys Gly Arg Lys Arg		705
705	710	715
Lys Gln Lys Leu Ile Ser Glu Glu Asp Leu		720
	725	730